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<u>L8</u>	L7 same (model or template or exemplar or prototype or reference)	10	<u>L8</u>
<u>L9</u>	L8 and (three near1 dimension\$3)	1	<u>L9</u>

<u>L10</u>	(dna or rna or chromoso\$5) near3 (analyz\$3 or analysis)	2	24875	<u>L10</u>
<u>L11</u>	L10 same image		690	<u>L11</u>
<u>L12</u>	L11 same (three near1 dimension\$3)		10	<u>L12</u>
<u>L13</u>	dna near2 image		363	<u>L13</u>
<u>L14</u>	L13 same densit\$3		16	<u>L14</u>
<u>L15</u>	L14 same ((two near1 dimension\$3) or (three near1 dimension\$3))		0	<u>L15</u>
<u>L16</u>	L14 and (two near1 dimension\$3)		4	<u>L16</u>
<u>L17</u>	L16 and (three near1 dimension\$3)		0	<u>L17</u>
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DB=PGPB; PLUR=YES; OP=ADJ		
<u>L3</u> dna same bit\$7 same model same image same (three nea	r2 dimension\$3)	0 <u>L3</u>
<u>L4</u> dna same 4model same image same (three near2 dimensi	ion\$3)	0 <u>L4</u>
<u>L5</u> dna same model same image same (three near2 dimension	on\$3)	5 <u>L5</u>
DB=USPT,EPAB,DWPI; PLUR=YES; OP=ADJ		
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L8 L7 same (model or template or exemplar or prototype or	reference) 1	0 <u>L8</u>
<u>L9</u> L8 and (three near1 dimension\$3)	'	1 <u>L9</u>

<u>L10</u>	(dna or rna or chromoso\$5) near3 (analyz\$3 or analysis)	24875	<u>L10</u>
<u>L11</u>	L10 same image	690	<u>L11</u>
<u>L12</u>	L11 same (three near1 dimension\$3)	10	<u>L12</u>
<u>L13</u>	dna near2 image	363	<u>L13</u>
<u>L14</u>	L13 same densit\$3	16	<u>L14</u>
<u>L15</u>	L14 same ((two near1 dimension\$3) or (three near1 dimension\$3))	0	<u>L15</u>
<u>L16</u>	L14 and (two near1 dimension\$3)	4	<u>L16</u>
<u>L17</u>	L16 and (three near1 dimension\$3)	0	<u>L17</u>
<u>L18</u>	L14 and (3d or (three near1 dimension\$3))	4	<u>L18</u>
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... DNA sequence occurs. The enzyme most commonly used for forensic DNA analysis is HaelII, which cuts DNA at the sequence 5'-GGCC-3'. ... www.biology.arizona.edu/human_bio/ problem_sets/DNA_forensics_2/03t.html - 8k - Cached - Similar pages

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DATE-ISSUED: June 29, 2004

INVENTOR-INFORMATION:

NAME

CITY

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DOCUMENT-IDENTIFIER: US 6445820 B1

TITLE: Method for conducting analysis of handwriting

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